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Novel human protei
Human polypeptide
Human peptide #94
Peptide #98 encode
Protein #88 encode
Human brain expres
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human bone marrow
Peptide #95 encode
Peptide #97 encode
Peptide #93 encode
                                                                     April 1, 2003, 08:45:42 ; Search time 75 Seconds (without alignments) 797.727 Million cell updates/sec
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Pred. No. is the number of results predicted by chance to have a score greater than or equal to the score of the result being printed, and is derived by analysis of the total score distribution.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                               1 MDRVYBIPEEPNVDPVSSLE........RTRVENSEPPFETEARQSVV 449
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Description
                                                                                                                                                                                                                                                                                                                                                                 GenCore version 5.1.4 p5_4578
Copyright (c) 1993 - 2003 Compugen Ltd.
                                                                                                                                                                                                                                       Total number of hits satisfying chosen parameters:
                                                                                                                                                                                                                908470 segs, 133250620 residues
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                                                                                                                                                                                                                                                                                                      Post-processing: Minimum Match 0%
Maximum Match 100%
Listing first 45 summaries
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ABB32592
ABB18089
AAM53421
AAM65799
AAM13661
AAM26060
                                                OM protein - protein search, using sw model
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ABB89300
                                                                                                                                                                            BLOSUM62
Gapop 10.0 , Gapext 0.5
                                                                                                                                                                                                                                                                                                                                                        A Geneseq 101002:*
. /SIDS2/gcgdata/
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Maximum DB seq length: 200000000
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2316
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Match Length
                                                                                                                          Title:
Perfect score:
                                                                                                                                                                            Scoring table:
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653
607
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                                                                         Run on:
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No.
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11 607 13 13 14 15 17 17 17 17 17 17 17 17 17 17	26.2 15.3 15.3 7.5	216 125 128 128	55553	ABG35433 AAU86530 ABB22596 AAM58002	Human peptide enco Novel human connec Protein #4595 enco
	4557	128 128	2222	AAU86530 ABB22596 AAM58002	Novel human connec Protein #4595 enco
	2 2 2 7	128	555	ABB22596 AAM58002	Procein #4595 enco
	ひてて	877	7 6	AAMSBU02	
	٠. ر. د. د.		•		Human brain expres
	۲.	86	7 (ABB29870	Peptide #2521 enco
		9 0	7 6	ABB35054	Peptide #2560 enco
	 	9 9	7 (ABB20466	Frocein #2465 enco
	, r	0 0	4 0	AAMARO40	Human bone marrow
	.5	98	22	AAM16062	Peptide #2496 enco
	7.5	98	22	AAM28558	Peptide #2595 enco
	7.5	98	22	AAM03789	Peptide #2471 enco
	7.5	986	53	ABG37776	Human peptide enco
	7.1	129	5	AAW81517	XK related Y (XKK)
		264	2 5	ABB91662 ABE03438	Herbicidally activ
		264	2 6	AAE03438 ARG63337	Human albumin fusi
	4.	989	5 5	AAU29285	Human PRO polypept
	6.4	785	23	ABB91027	Herbicidally activ
	4.8	264	22	AAE03484	Human gene 12 enco
	4.	264	53	ABG63335	Human albumin fusi
	9.	497	2 20	AAW95461	L. nelveticus pept
	5. 4 O A	457	2 6	ABF335/5	Staphylococcus epi
	1, 4 D n	1004	2 6	AAM99300	Salmonella typhi
	i d	786	3 6	AAU36408 AAI34510	E coli cellular r
	i 4	8 2 2	3 6	ABP27959	Streptococcus poly
	. 4.	353	22	ABG07351	Novel human diagno
	4.3	800	23	ABP40810	lococcu
	4.3	663	22	AAB85767	Human seven-transm
	4.3	408	20	AAY19788	B. burgdorferi ant
	4. E.	440	200	AAY19787	B. burgdorferi ant
	4. d	# T	2 6	AAWUYUZU	Timen cholinordio
44 45 99	4. 4. J. W.	1299	77	ABB70025	Drosophila melanog
				ALIGNMENTS	
RESULT 1					
97282	standard; Protein;	Prote		410 AA.	
XX AC ABB97282;					
DT 27-JUN-2002	(first	t entry)	ک		
DE Novel human	protein	n SEQ ID	ON GI): 550.	
		•		:	
Human; antiinf	н —	<pre>, vulnerary; erebroprotec</pre>	erar prot	11ntlammatory; ; cytostatic; x	<pre>immunomodulator; rheumatic; gene therapy;</pre>
KW neuroprotect KW expressed se	υ	antiparkinsonian; e tag.	kins	protein thera	EST;
XX OS Homo sapiens	· m				
XX PN WO200222660-122	242				
	:				

Ren F;

Zhao QA,

Tang YT, Liu C, Zhou P, Asundi V, Zhang J, Xue AJ, Yang Y, Wehrman T, Drmanac RT;

(HYSE-) HYSEQ INC.

WPI; 2002-292408/33. N-PSDB; ABN32468.

10-SEP-2001; 2001WO-US26015. 11-SEP-2000; 2000US-0659671.

21-MAR-2002.

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                                                                                      The present invention provides the protein and coding sequences of 444 novel human proteins. These were isolated from expressed sequences tags (ESTB). They can be used to stimulate cell growth, to regulate haematopolesis e.g. to treat aplastic anaema, to help tissue regrowth e.g. in burn treatment, to regulate the immune system e.g. to treat multiple sclerosis, to regulate activin or inhibin e.g. to treat infertility, to regulate haemostasis or thrombolysis e.g. to treat stroke and cancer, to screen for drugs, to treat inflammatory conditions e.g. rheumatodid arthritis, and to treat nervous system disorders e.g. Parkinson's disease. The present sequence is a protein of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     243 TIVLCFLTLLYTGINMPCWSAVQLKIDSPDLISKSHNWYQLLVYYMIRFIENAILLLLWY 302
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           YVSLISAEVPLGRVVLMVFSLVSVTYGATLCNMLAIQIKYDDYKIRLGPLEVLCITIWRT 248
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               LEITSRLLILUVLFSATLKLKAVPFLVLNFLIILFEPWIKFWRSGAOMPNNIEKNFSRVGT 308
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            183 FEIATRUVULVLFTSVLKTWVVIILINFSFFLYPWILFWCSGSPFPENIEKALSRVGT 242
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              309 LVVLISVTILYAGINFSCWSALQLRLADRDLVDKGQNWGHMGLHYSVRLVENVIMVLVPK 368
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            TFSFFMFSSIMVQLTLIFVHRDLAKDKPLSLFWHLILLGPVIRCLEAMIKYLTLWKKEEQ 129
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       369 FFGVKVLLNYCHSLIALQLIIAYLISIDFMLLFFQYLHPLRSLFTHNVVD----YLHCVC 424
An isolated polynucleotide for treating diseases associated with its encoded polypeptide such as cancer and multiple sclerosis -
                                                                                                                                                                                                                                                                                                                                                                                                                                 6 TLLFSLLPCALVQLTLLFVHRDLSRDRPLVLLLHLLQLGPLFRCFEVFCIY---FQSGNN 62
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      EEPYVSLTRKK-MLIDGEEVLIEWEVGHSIRTLAMHRNAYKRMSQIQAFLGSVPQLTYQL
                                                                                                                                                                                                                                                                                                                                                                         10;
                                                                                                                                                                                                                                                                                                                                Length 410;
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Similarity 45.8%; Pred. No. 8.8e-93;
57; Conservative 75; Mismatches 113; Indels
                                                       Example 2; SEQ ID NO 550; 509pp; English
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                                                                                                                                                                                                                                                                                                  410 AA;
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                                                                                                                                                                                                                                                                                                                                                                         Matches 167;
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                                                                                                                                                                                                                                                                                                Sequence
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The invention relates to novel genes (ABL89449-ABL90853) and proteins (ABB89040-ABB90444) useful for preventing, treating or ameliorating medical conditions e.g. by protein or gene therapy. The genes are isolated from a range of human tissues disclosed in the specification. The nucleic acids, proteins, antibodies and (ant) agonists are useful in the diagnosis, treatment and prevention of: (a) cancer, e.g. breast and ovarian cancer and other cancers of the adrenal gland, bone marrow, breast, gastrointestinal tract, liver, lung, or urogenital; (b) immune disorders e.g. Addison's disease, allergies, autoimmune themolytic anaemia, autoimmune thyroiditis, diabetes mellitus, Crohn's disease, multiple sclerosis, rhoumatoid arthritis and ulcerative colitis; (c) cardiovascular disorders such as myocardial ischaemias; (d) wound healing; (e) neurological diseases e.g. cerebral anoxia and parastic infections.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Human peptide #94 encoded by breast cell single exon nucleic acid probe.
                                                                                                                                                                                       Novel 1405 isolated polypeptides, useful for diagnosis, treatment and prevention of neural, immune system, muscular, reproductive, gastrointestinal, pulmonary, cardiovascular, renal and proliferative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              61 VKLVENVIMVLVFKFXGVKVXLNYCHXLXALQLIIAYLISIGFMLLFFQYLHPLRSLFTH 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     295 MPNNIEKNFSRVGTLVVLISVTILYAGINFSCWSALQIRLADRDLVDKGQNWGHMGLHYS 354
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            355 VRLVENVIMVLVFKFFGVKVLLNYCHSLIALQLIIAYLISIDFMLLFFQYLHPLRSLFTH
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                                                                                                                                                                                                                                                                                  Claim 11; SEQ ID NO 1676; 2081pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Length 131;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Score 653; DB 23;
Pred. No. 5.5e-68;
1; Mismatches 6;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             ABB27443 standard; Peptide; 216 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             28.2%;
18-MAY-2001; 2001WO-US16450.
                                 19-MAY-2000; 2000US-205515P.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Conservative
                                                                    (HUMA-) HUMAN GENOME
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    NVVDYLHCVCC 425
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       121 NVVDYLHCVCC 131
                                                                                                     Rosen CA;
                                                                                                                                       WPI; 2002-122018/16.
N-PSDB; ABL89709.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 131 AA;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            disease; cancer.
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                                                                                                                                                                                                                                               disorders
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Peptide #98 encoded by human foetal liver single exon nucleic acid probe.
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Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from NIPO at ftp.wipo.int/pub/published_pct_sequences.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                273 LVLNFLIILFEPWIKFWRSGAQMPNNIEKNFSRVGTLVVLISVTILYAGINFSCWSALQL 332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     333 RLADRDLVDKGQNWGHMGLHYSVRLVENVIMVLVFKFFGVKVLLNYCHSLIALQLIIAYL 392
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Gaps
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                                                                                      Human; foetal liver; gene expression; single exon nucleic acid probe
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              1 TYGAIRCNILAIQISNDDTTIKLPPIEFFCVVMMRFLEVISRVVTLAFFIASLKLKSLPV
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Human genome-derived single exon nucleic acid probes useful for analyzing gene expression in human fetal liver -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                47; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   26.2%; Score 607; DB 22; 55.1%; Pred. No. 2.9e-62;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              42; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Rank DR:
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       (MOLE-) MOLECULAR DYNAMICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chen W,
                                                                                                                                                                                                                                                                                                       04-FEB-2000, 2000US-0180312.
26-MAY-2000, 2000US-0207456.
30-UJN-2000; 2000US-0608408.
03-AUG-2000; 2000US-053366.
21-SEP-2000; 2000US-0234687.
27-SEP-2000; 2000US-0234687.
04-OCT-2000; 2000GB-0024263.
                                                                                                                                                                                                                                                             30-JAN-2001; 2001WO-US00669
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LATGFMLLFYQYLYPWQS 198
  (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Penn SG, Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              WPI; 2001-483447/52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              216 AA;
  01-FEB-2002
                                                                                                                                  Homo sapiens
                                                                                                                                                                                                                         39-AUG-2001
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Sequence
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ABB18089
ID ABB1
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    curcleic acid probes for measuring gene expression in a sample derived from whuman breast and BT 474 cells. The method involves contacting the probes with a collection of detectably labelled nucleic acids derived from mRNA of human breast, and then measuring the label.

C derived from mRNA of human breast, and then measuring the label.

C verifying the expression of regions of genomic DNA predicted to encode proteins. They are useful for gene discovery, and for determining predisposition and/or prognosing breast disease. Gene expression analysis is useful for assessing the toxicity of chemical agents on cells. The microarray of this invention presents a far greater diversity of probes for measuring gene expression, with far less bias than expressed sequence tag microarrays. The method is suitable for rapid production of functional information from genomic sequence. The present sequence is a peptide encoded by a single exon nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ö
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  probe of the invention.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          273 LVLNFLIILFEPWIKFWRSGAQMPNNIEKNFSRVGTLVVLISVTILYAGINFSCWSALQL 332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                61 LLIIYFVSLLAPWLEFWKSGAHLPGNKENNSNMVGTVLMLFLITLLYAAINFSCWSAVKL 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       213 TYGATLCNMLAIQIKYDDYKIRLGPLEVLCITIWRTLEITSRLLILVLFSATLKLKAVPF 272
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              The invention relates to a spatially-addressable set of single exon
                                                                                                                                                                                                                                                                                                                                                                                              New spatially-addressable set of single exon nucleic acid probes, useful for measuring gene expression in sample derived from human breast, comprises number of single exon nucleic acid probes -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         ;
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 27; SEQ ID NO 10411; 327pp + sequence listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         26.2%; Score 607; DB 22; Length 216; 55.1%; Pred. No. 2.9e-62; Live 42; Mismatches 47; Indels (
                                                                                                                                                                                                                                                                                                           Rank DR
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            ABB32592 standard; Peptide; 216 AA
                                                                                                                                                                                                                                                                MOLE-) MOLECULAR DYNAMICS INC
                                                                                                                                                                                                                                                                                                           Chen W,
                                                                                                                             2000US-0608408.
2000US-0632366.
2000US-0234687.
2000US-0236359.
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                                         30-JAN-2001; 2001WO-US00662
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181 LATGFMLLFYQYLYPWQS 198
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Matches 109; Conservative
                                                                                                                                                                                                                                                                                                           Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                  WPI; 2001-496933/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   216 AA
                                                                                                                                                  03-AUG-2000;
21-SEP-2000;
                                                                                                                                                                                                27-SEP-2000;
                                                                                                                                  30-JUN-2000;
                                                                                                           26-MAY-2000
09-AUG-2001
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                                                                                                                                                                                                                                                                                                           Penn SG,
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ABB32592
ID ABB3
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AC ABB3
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05-NOV-2001

AAM53421;

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measuring human gene expression in a sample derived from human heart (see PBA21355-ABA41305). The present sequence is a protein encoded by one such probe. The probes may be used for predicting, measuring and displaying gene expression in samples derived from the human heart via microarrays. By measuring gene expression, the probes are useful for predicting. dispnosing, grading, staging, monitoring and prognosing diseases of the human heart and vascular system e.g. cardiovascular disease. Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                          for analyzing gene expression in human
              Protein #88 encoded by probe for measuring heart cell gene expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  61 LLIIYFVSLLAPWLBFWKSGAHLPGNKENNSNNVGTVLMLFLITLLXAAINFSCWSAVKL 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              213 TYGATLCNMLAIQIKYDDYKIRLGPLEVLCITIWRTLEITSRLLILVLFSATLKLKAVPF 272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          present invention relates to single exon nucleic acid probes for
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   LVLNFLIILFEPWIKFWRSGAQMPNNIEKNFSRVGTLVVLISVTILYAGINFSCWSALQL
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Length 216;
                                           gene expression; heart; microarray; vascular syster
vascular disease; hypertension; cardiac arrhythmia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           47;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Query Match 26.2%; Score 607; DB 22; Best Local Similarity 55.1%; Pred. No. 2.9e-62; Matches 109; Conservative 42; Mismatches 47;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Claim 15; SEQ ID No 19859; 530pp; English.
                                                                                                                                                                                                                                                                                                                                                                                B.
                                                                                                                                                                                                                                                                                                                                                                                 Rank
                                                                                                                                                                                                                                                                                                                                                                                                                                          Single exon nucleic acid probes hearts -
                                                                                                                                                                                                                                                                                                                                               (MOLE-) MOLECULAR DYNAMICS INC
                                                                                                                                                                                                                                                                                                                                                                                Chen W,
                                                                                                                                                                                                                                          2000US-0207456,
2000US-0608408
                                                                                                                                                                                                                                                                   03-AUG-2000; 2000US-0632366.
21-SEP-2000; 2000US-0234687.
27-SEP-2000; 2000US-0236359.
04-OCT-2000; 2000GB-0024263.
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                                                                                                                                                                                                                            2000US-0180312
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                                                                                                                                                                                                                                                                                                                                                                                Hanzel DK,
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                                                           cardiovascular
                                                                                                                                   WO200157274-A2
                                                                                                        Homo sapiens
                                                                                                                                                                                                                                          26-MAY-2000;
                                                                                                                                                                                                                          04-FEB-2000;
                                                                                                                                                                09-AUG-2001
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                                                                                                                                                                                                                                                                                                                                                                             SG,
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The present invention provides a number of single exon nucleic acid probes which are derived from genomic sequences expressed in the human brain. They can be used to measure gene expression in brain cell samples, which may enable the diagnosis and improved treatment of nervous system diseases such as Alzheimer's disease, multiple sclerosis, schizophrenia, epilepsy and cancers. The present sequence is a protein encoded by one of the probes of the invention.
                                                                   Human brain expressed single exon probe encoded protein SEQ ID NO: 25526.
                                                                                                                                                                                                                                                                                                                                                                                                                                                             Single exon nucleic acid probes for analyzing gene expression in human
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             213 TYGATLCNMLAIQIKYDDYKIRLGPLEVLCITIWRTLEITSRLLILVLFSATLKLKAVPF 272
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Human; brain expressed exon; gene expression analysis; probe; microarray; Alzheimer's disease; multiple sclerosis; schizophrenia.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    47;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         DB 22;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       26.2%; Score 607; DB 22;
55.1%; Pred. No. 2.9e-62;
ive 42; Mismatches 47
                                                                                                                                                                                                                                                                                                                                                                                                        DR;
                                                                                                                                                                                                                                                                                                                                                                                                        Rank
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AAM65799
ID AAM65799 standard; Protein; 216 AA.
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                                                                                                                                                                                                                                                                                                                                                                            (MOLE-) MOLECULAR DYNAMICS INC
                                                                                                                                                                                                                                                                                                                                                                                                        Chen W,
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21-SEP-2000; 2000US-0234687.
27-SEP-2000; 2000US-0236359.
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2000US-0608408.
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                                                                                                                                                                                                                                                                                                                                                                                                        Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                   WPI; 2001-483446/52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Local Similarity
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                                                                                                                         epilepsy; cancer
                                                                                                                                                                               WO200157275-A2
                                                                                                                                                                                                                                                               04-FEB-2000;
26-MAY-2000;
30-JUN-2000;
03-AUG-2000;
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AAM53421 standard; Protein; 216 AA.

RESULT 6 AAM53421 ID AAM S

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Peptide #95 encoded by probe for measuring cervical gene expression.
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                        (MOLE-) MOLECULAR DYNAMICS INC
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Chen W,
                                                                                                                                                                                                                                                                                              04-FEB-2000; 2000US-0180312.
26-MAY-2000; 2000US-0207456.
30-UUN-2000; 2000US-0608408.
03-AUG-2000; 2000US-0633366.
21-SEP-2000; 2000US-0236559.
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LATGFMLLFYQYLYPWQS 198
                                                                                                                                                                                                                                                                                                                                                                                                                              04-OCT-2000; 2000GB-0024263
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                                                                                        cervical cancer
                                                                                                                                                                       WO200157278-A2
                                                                                                                               Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           probes which are derived from genomic sequences expressed in the human bone marrow. They can be used to measure gene expression in bone marrow samples, which may enable the improved diagnosis and treatment of cancers such as lymphoma, leukaemia and myeloma. The present sequence is a protein encoded by one of the probes of the invention.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         213 TYGATLCNMLAIQIKYDDYKIRLGPLEVLCITIWRTLEITSRLLILVLFSATLKLKAVPF 272
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          273 LVLNFLIILFEPWIKFWRSGAOMPNNIEKNFSRVGTLVVLISVTILYAGINFSCWSALOL 332
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     61 LLIYFVSLLAPWLEFWKSGAHLPGNKENNSNWVGTVLMLFLITLLYAAINFSCWSAVKL 120
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              RLADRDLVDKGQNWGHMGLHYSVRLVENVIMVLVFKFFGVKVLLNYCHSLIALQLIIAYL 392
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  0; Gaps
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         provides a number of single exon nucleic acid
                                                                                                                             Human; bone marrow expressed exon; gene expression analysis; probe;
microarray; cancer; leukaemia; lymphoma; myeloma.
                                                                                 Human bone marrow expressed probe encoded protein SEQ ID NO: 26105
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Example 4; SEQ ID NO: 26105; 658pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 26.2%; Score 607; DB 22; Length 216; 55.1%; Pred. No. 2.9e-62; tive 42; Mismatches 47; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  Human genome-derived single exon nucleic acid probes useful analyzing gene expression in human bone marrow -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Chen W, Rank DR
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      (MOLE-) MOLECULAR DYNAMICS INC
                                                                                                                           expressed
                                                                                                                                                                                                                                                                                                                                                                                ; 2000US-0207456.
; 2000US-0608408.
; 2000US-0632366.
; 2000US-0234687.
                                                                                                                                                                                                                                                                                                                    30-JAN-2001; 2001WO-US00668
                                                                                                                                                                                                                                                                                                                                                                                                                                                                      27-SEP-2000; 2000US-0236359 04-OCT-2000; 2000GB-0024263
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LATGFMLLFYQYLYPWQS 198
                                       (first entry)
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        WPI; 2001-488900/53
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 Query Match
Best Local Similarity
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                                                                                                                                                                                                                                 WO200157276-A2
                                                                                                                                                                                           Homo sapiens
                                                                                                                                                                                                                                                                                                                                                                                                    30-JUN-2000;
                                                                                                                                                                                                                                                                                                                                                                                                                                              21-SEP-2000;
                                                                                                                                                                                                                                                                                                                                                                                    26-MAY-2000;
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AAM65799;
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The present invention relates to human single exon nucleic acid probes (SENP: see AAI10068-AAI28459). The present sequence is a peptide encoded by one such probe. The SENPs are derived from human HeLa cells. The SENPs and be used to produce a single exon microarray, which can be used for measuring human gene expression in a sample derived from human cervical epithelial cells. By measuring gene expression, the probes are therefore useful in grading and/or staging of diseases of the cervix, notably
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               the printed
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Note: The sequence data for this patent did not form part of the printer specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
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human; microarray; gene expression; cervical epithelial cell;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                  ô
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Human genome-derived single exon nucleic acid probes useful f
analyzing gene expression in human cervical epithelial cells
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 42; Mismatches
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Claim 27; SEQ ID No 18487; 487pp; English
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entry)

(first

17-OCT-2001

Peptide #97

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The present invention relates to novel single exon nucleic acid probes (see AA100010-AA110067). The present sequence is a peptide encoded by one such probes. The probes are useful for measuring human gene expression in a human breast sample, where the probe hybridises at high stringency to ancleic acid expressed in the human breast. The probes are useful for predicting, diagnosing, grading, staging, monitoring and prognosing diseases of the human breast, particularly those diseases with polygenic aetiology. The diseases include: breast cancer, disorders of development, inflammatory diseases of the breast, fibrocystic changes, proliferative
                                    Probe; human; breast disease; breast cancer; development disorder; inflammatory disease; proliferative breast disease; non-carcinoma tumour.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              breast disease and non-carcinoma tumours.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
Peptide #93 encoded by probe for measuring human breast gene expression.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Novel single exon nucleic acid probe used to measuring gene expression in a human breast \cdot
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 213 TYGATLCHMLAIQIKYDDYKIRLGPLEVLCITIWRTLEITSRLLILVLFSATLKLKAVPF 272
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               Claim 27; SEQ ID No 10151; 322pp; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                      DR;
                                                                                                                                                                                                                                                                                                                                                                                                                                                      Rank
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                                                                                                                                                                                                                                                                                                                                                                                                                                                      Chen W,
                                                                                                                                                                                                                     29-JAN-2001; 2001WO-US00661
                                                                                                                                                                                                                                                                              2000US-0207456
2000US-0608408
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27-SEP-2000; 2000US-0236359.
04-OCT-2000; 2000GB-0024263.
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LATGFMLLFYQYLYPWQS
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                                                                                                                                          WO200157270-A2
                                                                                                                                                                                                                                                                              26-MAY-2000;
                                                                                                   Homo sapiens
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     LVLNFLIILFEPWIKFWRSGAQMPNNIEKNFSRVGTLVVLISVTILYAGINFSCWSALQL 332
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                                                            encoded by probe for measuring placental gene expression
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 single exon nucleic acid probes useful for
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                                                                                                   Probe; microarray; human; placenta; antenatal diagnosis; genetic disorder.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     analyzing gene expression in human placenta
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       Claim 27; SEQ ID No 26329; 654pp; English.
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(MOLE-) MOLECULAR DYNAMICS INC

04-OCT-2000;

Hanzel DK,

Penn SG,

Human genome-derived

WPI; 2001-488897/53

2000US-0207456. 2000US-0608408. 2000US-0632366. 2000US-0234687. 2000US-0234687.

26-MAY-2000; 03-AUG-2000; 21-SEP-2000;

34-FEB-2000;

30-JAN-2001; 2001WO-US00663.

WO200157272-A2.

39-AUG-2001

Homo sapiens

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Local Simi hes 109;

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Sequence Query Match (first entry)

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AAM01411 AAM01411;

AAM01411 ID AAM(XX AC AAM(XX DT 09-C

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Gaps

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47; Indels

Length 216;

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WPI, 2002-114183/15.
                                  W0200186003-A2
                              Homo sapiens
                                                 10-JUN-2000;
                                                     21-SEP-2000;
                                             04-FEB-2000;
                                                   33-AUG-2000;
   19-AUG-2002
                                               26-MAY-2000
                                      15-NOV-2001
ABG35433;
                                                                Penn SG,
                                                             (MOLE-)
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haemosiderosis, pulmonary histiocytosis, lymphangioleiomyomtosis, pulmonary alveolar proteinosis, Karagener syndrome, fibrocystic pulmonary dysplasia, primary ciliary dyskinesis, pulmonary hypertension and hyaline membrane disease. The present sequence is a peptide/protein encoded by a single exon probe of the invention.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at they wipo.int/pub/published_pot_sequences.
Niemann-Pick disease, Hermansky-Pudlak syndrome, sarcoidosis, pulmonary
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   RLADRDLVDKGQNWGHMGLHYSVRLVENVIMVLVFKFFGVKVLLNYCHSLIALQLIIAYL 392
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Human; connective tissue related disorder; cancer; cytostatic.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Novel human connective tissue related polypeptide #96.
                                                                                                                                                                                                                                                                                                                                                               26.2%; Score 607; DB 23; 55.1%; Pred. No. 2.9e-62; ive 42; Mismatches 47;
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2000US-0184664.
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2000US-0205515.
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2000US-0215135.
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181 LATGFMLLFYQYLYPWQS 198
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Matches 109; Conservative
                                                                                                                                                                                                                                                                                                                216 AA;
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                                                                                                             Human peptide encoded by genome-derived single exon probe SEQ ID 25098
                                                                                                                                             Human; single exon probe; asthma; lung cancer; COPD; ILD; chronic obstructive pulmonary disease; interstitial lung disease; familial idiopathic pulmonary fibrosis; neurofibromatosis; tuberous sclerosis; Gaucher's disease; Hermansky-Pudlak syndrome; sarcoidosis; pulmonary haemosiderosis; pulmonary hietlocytosis; lymphangioleiomyomtosis; Karagener syndrome; pulmonary alveolar proteinosis; fibrocystic pulmonary dyskinesis; pulmonary alveolar proteinosis; fibrocystic pulmonary dysplasia; primary ciliary dyskinesis; pulmonary hypertension;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                            Spatially-addressable set of single exon nucleic acid probes, used to
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         measure gene expression in human lung samples
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           Claim 27; SEQ ID No 25098; 634pp; English.
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2000US-0632366.
2000US-234687P.
2000US-236359P.
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2000US-0227182.
2000US-0227009.
2000US-0228924.
                  2000US-0225758
2000US-0225759
                      2000US-0226279
2000US-0226681
                                          0000S-0229509
                              23-AUG-2000;
30-AUG-2000;
01-SEP-2000;
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The present invention relates to the isolation of novel human connective tissue related polypeptides and the polynucleotide (CDNA and genomic) sequences encoding them. The sequences of the invention are useful in the diagnosis, treatment, prevention and/or prognosis of diseases associated with connective tissue(s), including cancer. The polynucleotide sequences of the invention are also useful in gene therapy, AAU86435-AAU86923 represent the novel human connective tissue related polypeptides.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Nucleic acid encoding novel connective tissue associated polypeptides, used in diagnosing, preventing, treating or ameliorating a disorder such as cancer or rheumatoid arthritis -
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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                Barash SC,
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N-PSDB; ABK41708.
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62 IMILVFRFFGGKTLLNCCDSLIAVQLIISYLLATGFMLLFYQYLYPWQS 110

362 IMVLVFKFFGVKVLLNYCHSLIALQLIIAYLISIDFMLLFFQYLHPLRS

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                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                             Protein #4595 encoded by probe for measuring heart cell gene expression.
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                                                                                             LISVTILYAGINFSCWSALQLRLADRDLVDKGQNWGHMGLHYSVRLVENVIMVLVFKFFG 371
                                        Gaps
                                                                                                                                  The present invention relates to single exon nucleic acid probes for
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د
   Pred. No. 1.6e-58;
                                     1; Mismatches
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2000US-0234687.
2000US-0236359.
94.78;
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04-OCT-2000; 2000GB-0024263.
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                            Matches 108; Conservative
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   Best Local Similarity
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probes which are derived from genomic sequences expressed in the human brain. They can be used to measure gene expression in brain cell samples, which may enable the diagnosis and improved treatment of nervous system diseases such as Alzheimer's disease, multiple sclerosis, schizophrenia, epilepsy and cancers. The present sequence is a protein encoded by one of the probes of the invention.
                                                                                                                  Human brain expressed single exon probe encoded protein SEQ ID NO: 30107.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        Single exon nucleic acid probes for analyzing gene expression in human brains -
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              Gaps
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         302 NFSRVGTLVVLISVTILYAGINPSCWSALQLRLADRDLVDKGQNWGHMGLHYSVRLVENV 361
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          The present invention provides a number of single exon nucleic acid
                                                                                                                                            Human; brain expressed exon; gene expression analysis; probe; microarray; Alzheimer's disease; multiple sclerosis; schizophrenia;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              .
0
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         Example 4; SEQ ID NO: 30107; 650pp + Sequence Listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   362 IMVLVFKFFGVKVLLNYCHSLIALQLIIAYLISIDFMLLFFQYLHPLRS 410
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              21; Indels
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                15.3%; Score 354; DB 22; 59.6%; Pred. No. 5.7e-33; ive 23; Mismatches 21;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     ДR.;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Rank
                            AAM58002 standard; Protein; 128 AA
                                                                                                                                                                                                                                                                                                                                                                                                                                         (MOLE-) MOLECULAR DYNAMICS INC
                                                                                                                                                                                                                                                                                                                                    26-MAY-2000; 2000US-0207456.
30-JUN-2000; 2000US-0608408.
03-AUG-2000; 2000US-0632366.
21-SEP-2000; 2000US-0234687.
27-SEP-2000; 2000US-0234687.
04-OCT-2000; 2000GB-0024263.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Chen W,
                                                                                                                                                                                                                                                                                            30-JAN-2001; 2001WO-US00667
                                                                                    (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                65; Conservative
                                                                                                                                                                                                                                                                                                                                                                                                                                                                     Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                WPI; 2001-483446/52
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   Query Match
Best Local Similarity
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      128 AA;
                                                                                                                                                                           epilepsy; cancer.
                                                                                                                                                                                                                                   WO200157275-A2
                                                                                                                                                                                                        Homo sapiens.
                                                                                                                                                                                                                                                                                                                       04-FEB-2000;
                                                                                     05-NOV-2001
                                                                                                                                                                                                                                                                09-AUG-2001
                                                       AAM58002;
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RESULT 14
               AAM58002
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RESULT 15

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302 NFSRVGTLVVLISVTILYAGINFSCWSALQLRLADRDLVDKGQNWGHMGLHYSVRLVENV 361

Conservative

65;

Matches

8

Local Similarity

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rucleic acid probes for measuring gene expression in a sample derived from human breast and Br 474 cells. The method involves contacting the probes with a collection of detectably labelled nucleic acids derived from mRNA of human breast, and then measuring the label country of human breast, and then measuring the label country of the microarray. The probes are useful for verifying the expression of regions of genomic DNA predicted to encode proteins. They are useful for gene discovery, and for determining predisposition and/or prognosing breast disease. Gene determining predisposition and/or prognosing breast disease. Gene expression analysis is useful for assessing the toxicity of chemical agents on cells. The microarray of this invention presents a far greater diversity of probes for measuring gene expression, with far less bias than expressed sequence tag microarrays. The method is suitable for rapid production of functional information from genomic sequence. The present sequence is a peptide encoded by a single exon nucleic acid
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          probe of the invention.

Note: The sequence data for this patent did not form part of the printed specification, but was obtained in electronic format directly from WIPO at ftp.wipo.int/pub/published_pct_sequences.
                                                                                                                              Peptide #2521 encoded by breast cell single exon nucleic acid probe.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                           The invention relates to a spatially-addressable set of single exon
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                          New spatially-addressable set of single exon nucleic acid probes, useful for measuring gene expression in sample derived from human breast, comprises number of single exon nucleic acid probes -
                                                                                                                                                                     Human; microarray; single exon probe; gene expression; breast;
disease; cancer.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Claim 27; SEQ ID NO 12838; 327pp + sequence listing; English.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Chen W, Rank DR;
                   ABB29870 standard; Peptide; 86 AA.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                               (MOLE-) MOLECULAR DYNAMICS INC.
                                                                                                                                                                                                                                                                                                                                                                                               2000US-0207456.
2000US-0608408.
2000US-0632366.
2000US-0234687.
                                                                                                                                                                                                                                                                                                                                      30-JAN-2001; 2001WO-US00662
                                                                                                                                                                                                                                                                                                                                                                                                                                                                       27-SEP-2000; 2000US-0236359.
04-OCT-2000; 2000GB-0024263.
                                                                                           01-FEB-2002 (first entry)
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      Hanzel DK,
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         WPI; 2001-496933/54.
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                         86 AA;
                                                                                                                                                                                                                                                             WO200157271-A2
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03-AUG-2000;
21-SEP-2000;
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                                                                                                                                                                                                                                                                                                                                                                                                 26-MAY-2000;
                                                                                                                                                                                                                                                                                                   09-AUG-2001
                                                        ABB29870;
                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                    Penn SG,
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Search completed: April 1, 2003, 08:47:08 Job time : 77 secs

/ Match 7.5%; Score 174.5; DB 22; Length 86; Local Similarity 53.4%; Pred. No. 3.3e-12; Nes 39; Conservative 12; Mismatches 21; Indels 1 Query Match Best Loca Matches

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Gaps

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130 EEPYVSLTRKK-MLIDGEEVLIEWEVGHSIRTLAMHRNAYKRMSQIQAFLGSVPQLTYQL 188

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189 YVSLISAEVPLGR 201

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